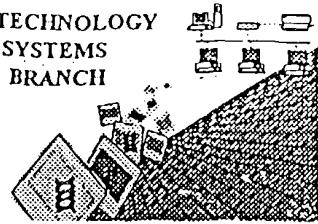


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 08/936,651A
Source: IFW/6
Date Processed by STIC: 3/29/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>08/936,657A</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input checked="" type="checkbox"/> Variable Length	Sequence(s) <u>12-13</u> contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) <u> </u> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) <u> </u> missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) <u> </u> missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <input type="checkbox"/> Use of <220>	Sequence(s) <u> </u> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <input type="checkbox"/> Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>	



IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/936,657A

DATE: 03/29/2004

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt
 Output Set: N:\CRF4\03292004\H936657A.raw

```

3 <110> APPLICANT: Sirna Therapeutics, Inc.
5 <120> TITLE OF INVENTION: Modified Ribozymes
7 <130> FILE REFERENCE: MBHB 00-838-A; 228/213
9 <140> CURRENT APPLICATION NUMBER: US 08/936657A
10 <141> CURRENT FILING DATE: 1997-09-24
12 <150> PRIOR APPLICATION NUMBER: US 07/965,411
13 <151> PRIOR FILING DATE: 1993-08-09
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: PatentIn version 3.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 32
21 <212> TYPE: DNA
22 <213> ORGANISM: Artificial Sequence
24 <220> FEATURE:
25 <223> OTHER INFORMATION: Ribozyme
27 <400> SEQUENCE: 1
28 gatattcctga ctccctatacg tgagtgcgtat ta
31 <210> SEQ ID NO: 2
32 <211> LENGTH: 36
33 <212> TYPE: DNA
34 <213> ORGANISM: Artificial Sequence
36 <220> FEATURE:
37 <223> OTHER INFORMATION: Ribozyme
39 <400> SEQUENCE: 2
40 taatacgtact cactataggg agtcaggata tctgca
43 <210> SEQ ID NO: 3
44 <211> LENGTH: 51
45 <212> TYPE: DNA
46 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
49 <223> OTHER INFORMATION: Ribozyme
51 <400> SEQUENCE: 3
52 ggagtttcgg cctaaccggcc tcatacgagg accctatagt gagtcgtatt a
55 <210> SEQ ID NO: 4
56 <211> LENGTH: 55
57 <212> TYPE: DNA
58 <213> ORGANISM: Artificial Sequence
60 <220> FEATURE:
61 <223> OTHER INFORMATION: Ribozyme
63 <400> SEQUENCE: 4
64 taatacgtact cactataggg tcctctgtatc aggccgttag gccgaaactc ctgca
67 <210> SEQ ID NO: 5
68 <211> LENGTH: 34

```

*Does Not Comply
Corrected Diskette Needed*

P.3

32

36

51

55

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/936,657A

DATE: 03/29/2004

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt
 Output Set: N:\CRF4\03292004\H936657A.raw

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69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: Ribozyme
75 <400> SEQUENCE: 5
76 ggguccucug augaggccgu uaggccgaaa cucc          34
79 <210> SEQ ID NO: 6
80 <211> LENGTH: 34
81 <212> TYPE: DNA
82 <213> ORGANISM: Artificial Sequence
84 <220> FEATURE:
85 <223> OTHER INFORMATION: Ribozyme
87 <220> FEATURE:
88 <221> NAME/KEY: misc_feature
89 <223> OTHER INFORMATION: All U nucleic acids are 2'-fluoro derivatives.
92 <400> SEQUENCE: 6
93 ggguccucug augaggccgu uaggccgaaa cucc          34
96 <210> SEQ ID NO: 7
97 <211> LENGTH: 34
98 <212> TYPE: DNA
99 <213> ORGANISM: Artificial Sequence
101 <220> FEATURE:
102 <223> OTHER INFORMATION: Ribozyme
104 <220> FEATURE:
105 <221> NAME/KEY: misc_feature
106 <223> OTHER INFORMATION: All U nucleic acids are 2'-amino derivatives.
109 <400> SEQUENCE: 7
110 ggguccucug augaggccgu uaggccgaaa cucc          34
113 <210> SEQ ID NO: 8
114 <211> LENGTH: 12
115 <212> TYPE: RNA
116 <213> ORGANISM: Artificial Sequence
118 <220> FEATURE:
119 <223> OTHER INFORMATION: Substrate
121 <400> SEQUENCE: 8
122 gggagucagg au                                12
125 <210> SEQ ID NO: 9
126 <211> LENGTH: 12
127 <212> TYPE: RNA
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Substrate
133 <220> FEATURE:
134 <221> NAME/KEY: misc_feature
135 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives
138 <400> SEQUENCE: 9
139 gggagucagg au                                12
142 <210> SEQ ID NO: 10
143 <211> LENGTH: 12

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RAW SEQUENCE LISTING

DATE: 03/29/2004

PATENT APPLICATION: US/08/936,657A

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

144 <212> TYPE: RNA
 145 <213> ORGANISM: Artificial Sequence
 147 <220> FEATURE:
 148 <223> OTHER INFORMATION: Substrate
 150 <220> FEATURE:
 151 <221> NAME/KEY: misc_feature
 152 <222> LOCATION: (7)..(7)
 153 <223> OTHER INFORMATION: 2'-fluoro derivative
 156 <400> SEQUENCE: 10
 157 gggagucagg au 12
 160 <210> SEQ ID NO: 11
 161 <211> LENGTH: 12
 162 <212> TYPE: RNA
 163 <213> ORGANISM: Artificial Sequence
 165 <220> FEATURE:
 166 <223> OTHER INFORMATION: Substrate
 168 <220> FEATURE:
 169 <221> NAME/KEY: misc_feature
 170 <222> LOCATION: (6)..(6)
 171 <223> OTHER INFORMATION: 2'-amino derivative
 174 <400> SEQUENCE: 11
 175 gggagucagg au 12
 178 <210> SEQ ID NO: 12
 179 <211> LENGTH: 13
 180 <212> TYPE: DNA
 181 <213> ORGANISM: Artificial Sequence
 183 <220> FEATURE:
 184 <223> OTHER INFORMATION: Substrate
 186 <220> FEATURE:
 187 <221> NAME/KEY: misc_feature
 188 <222> LOCATION: (13)..(13)
 189 <223> OTHER INFORMATION: N stands for any one or more A, T, G, C.
 192 <400> SEQUENCE: 12
 W--> 193 gggagucagg aun 13
 196 <210> SEQ ID NO: 13
 197 <211> LENGTH: 13
 198 <212> TYPE: DNA
 199 <213> ORGANISM: Artificial Sequence
 201 <220> FEATURE:
 202 <223> OTHER INFORMATION: Substrate
 204 <220> FEATURE:
 205 <221> NAME/KEY: misc_feature
 206 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives.
 209 <220> FEATURE:
 210 <221> NAME/KEY: misc_feature
 211 <222> LOCATION: (13)..(13)
 212 <223> OTHER INFORMATION: N stands for one or more modified or unmodified nucleotides.
 215 <400> SEQUENCE: 13
 W--> 216 gggagucagg aun 13

variable length is not permitted.
 see item 5
 on Error
 summary
 sheet.

same error

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/936,657A

DATE: 03/29/2004

TIME: 10:30:20

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

219 <210> SEQ ID NO: 14
220 <211> LENGTH: 12
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Substrate
227 <220> FEATURE:
228 <221> NAME/KEY: misc_feature
229 <223> OTHER INFORMATION: All U nucleotides are 2'-fluoro derivatives.
232 <400> SEQUENCE: 14
233 gggagucagg au 12
236 <210> SEQ ID NO: 15
237 <211> LENGTH: 36
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Ribozyme
244 <400> SEQUENCE: 15
245 cacaacacug augaggccgu uaggccgaaa cgggca 36

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/29/2004
PATENT APPLICATION: US/08/936,657A TIME: 10:30:21

Input Set : A:\00-838-A.SequenceListing.txt
Output Set: N:\CRF4\03292004\H936657A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:12; N Pos. 13

Seq#:13; N Pos. 13

VERIFICATION SUMMARY

PATENT APPLICATION: US/08/936,657A

DATE: 03/29/2004

TIME: 10:30:21

Input Set : A:\00-838-A.SequenceListing.txt

Output Set: N:\CRF4\03292004\H936657A.raw

L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0

L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0